

RAW SEQUENCE LISTING

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Application Serial Number: 10/578,701
Source: IFWP
Date Processed by STIC: 5/18/06

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IFWP

RAW SEQUENCE LISTING

DATE: 05/18/2006

PATENT APPLICATION: US/10/578,701

TIME: 10:27:53

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\05182006\J578701.raw

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3 <110> APPLICANT: BioVentures, Inc.
4     Vanderbilt University
5     Dawson, Elliot P.
6     Phillips, John A.
7     Womble, Kristie E.
9 <120> TITLE OF INVENTION: Chromosome 5 Genetic Variants Related to Dyslexia
11 <130> FILE REFERENCE: 14160-2US
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/578,701
C--> 13 <141> CURRENT FILING DATE: 2006-05-09
13 <150> PRIOR APPLICATION NUMBER: 60/520,366
14 <151> PRIOR FILING DATE: 2003-11-14
16 <150> PRIOR APPLICATION NUMBER: PCT/US04/37587
17 <151> PRIOR FILING DATE: 2004-11-13
19 <160> NUMBER OF SEQ ID NOS: 25
21 <170> SOFTWARE: PatentIn version 3.2
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 3664
25 <212> TYPE: DNA
26 <213> ORGANISM: Homo sapiens
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31 attcatcatg tgcatacaag gccattctgt gtttcctact cttgccttgg gctcatcatt      120
33 attaattctg aattccattt gttcttcact ttttgaatat gtctgtttag ttgactgtag      180
35 tgccactggc aggaccatgt gccaggaaa tccaagactc atatttggac gaaagctatg      240
37 tccacttttc aactagtacc cctacccaaa ttaccatagc aacacaaaaa ttgcagatgc      300
39 ctacattcta gaatcatgtt ctaaagggat gtcatcattt acaaaatgtc tttgttgagt      360
41 ctgaatgggt caaacaatag caaaaaagga ttatttctct cttggacatt tcaaagtact      420
43 atgacacaaa atatccaaga cttgttatgg tgaggagcca agtggaatgg aaaggacagc      480
45 tcatcccggc ggctgggagt gcatgcacac acatgccccc tttttcttgc ctactaacag      540
47 gatctataga aggcgtacat aatgagtatg taggggactt ggctgcttct agttaggaat      600
49 gagacactga tatggttggg atatagtaag agaaaaaggg aggtctttct taaaaaacgg      660
51 ttttgtgtaa aaatagagat ggcacttaat ggatatcata ttagcaggct ccctggacaa      720
53 atacatagag ccaaaacttc tcatcgatta gccacctctt caagtttagg ggttgaaaat      780
55 ctgaaacaac tacaaacatg gtatctctct gaaaaggaga taacgtaaaa gttatcacat      840
57 attaataata tgtgtatgaa taaattgaca agctggtagg aaattagaaa taaaagtctt      900
59 gaggcaataa aagaggtaat aacataggca aaaagagctc ttcttctgga gagtggttgt      960
61 agatggagta aacaagttta ggtactgaac tgagaatagc acatggatag accaattgtg     1020
63 gatgaaggag actaaagaga ggtttaacga atattgaaat gaacctccag gtaggttgta     1080
65 tttattagtt tgctgggaac aagctgcttt tctctctcct gtgaagcagg aaggcaaatt     1140
67 tctagtggct ttccaaagga aatgggaaat ctaaggaaat ggtttgatac cagagtgttc     1200
69 tccttagggt tattttaatg atggacttaa agatactttc ctatactcat gagctatggt     1260
71 gtctctgata ttctttggta tattttacca aaaagataga ataggtgcca caagtattaa     1320
73 aaattttaga ctctcagag cattacaaaa aacaagcaca aaatagaage ctaatatgca     1380

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75 gggaaagtca ctgaccatgc ccttggtact gctgattgta ttgcagagca agagatggac 1440
77 cctgagggtta cttgaagcca acaagtttca cttctggaaa aagacttcag aatatgagtt 1500
79 taaaatataa aaaggaatt tgagccaaga cacaagaaca aacttttttt gacaattata 1560
81 tctttattat tcctcttaca gagctacatt tactcttact aagtttcaga gtcaggtagt 1620
83 aatttacagt aagactgaat taccatccat aacgttagat gtccttattg aaacttcaac 1680
85 atcatttcca aatatcagca ttagcattgt gcttgacatt catttaacga agttactgaa 1740
87 aatctattaa gtataagaca tcagttattt ttaatagaag tttctgaaaa catttcagca 1800
89 aaatagcctg ttgagaaaaa tgtgtatgct gaaaaaaaaa aatgaacaaa taggaaagcc 1860
91 tgggttcacaa acaggtgtca gggaaataga cagtactttt atagtaataa cataagaaca 1920
93 aacttcttga aggtaagttt tattaaataa taggacaaca acaagataaa atgacttctt 1980
95 cctgatattt atatatgat tgctggctgg tcataagact gtttttaggc aacgtgtttt 2040
97 gaaaaaccag aaagtctact accttgagtt ttcagccacg tgagaatagc aagattcagt 2100
99 gtttatactt gatagcatct taattaggcc tacaggcctc cctttcacat aactaccttc 2160
101 aagtttatga cagctcaaac tcacaattat cattatggag aagagagaag agttaagcta 2220
103 aaaacagacc actttcagag gacctgaaag caacgtaatc agtcacctat tgccatatac 2280
105 aagccacccc caaacataat gacttaaaac agcgatcatc tattattgct tatgagtctc 2340
107 tgagtcagct gaacattcct gctgatctgg gcttggttag gcttatttta gctgtgttca 2400
109 ttcttggctc gcagatagct gacaatcacc taggggctga ctgtaggcat tccagctgag 2460
111 atatgctctc tgtgtctttt atccttttagc aggaggaggc ttgctcacag ggtggttaca 2520
113 ggcattccaag agagtcagca taaatgtgaa aagtttccaa aatatcagat tcagtcctat 2580
115 gtaatctggg ttccattgca ttctcttggc cagagcaagt tgcaagacaa gtccaaattc 2640
117 aagaagggtca agaaatacac tccattctca ggtaggagaa gctgcaaaga actgtgacaa 2700
119 tctatgacaa atagtatggt caaagggaat aatatgggaa gatgtgccct ccgccaactt 2760
121 ctgaggga aaatacagctt ttgtaattat tagtaatata gactgtctaa tatttctaga 2820
123 gaaatctatg actttgagtt gaaatatctg aggccaacac tccaagcaat tttaaacaag 2880
125 tgggtgacaga aattaccaga cacacatcaa gactcaagta taaagctata caatttaagg 2940
127 atgctcagca aatgttactg aattgactgg gtagtcctta aagagctgaa gaataaaaga 3000
129 tgttatgaga aatccaacaa taccaaatat aaattgcctc aggttctgaa atattcaata 3060
131 aagtattctc actgtagttc cttcagctta gctgatttgg actttggctg tgaaaacatt 3120
133 atcctcagtg tttaaaaggt tggaaaattc tactgggtct ttggcccaac ctggaattaa 3180
135 atcctgatgc ttagaacctc aaagtctaaa atcttctatt gtcactttac agagctattg 3240
137 aaacatatta ataaacttgt atcatactga tttgattcta atttttgtgg gacattgttt 3300
139 aaaaattgtt gaaatgcata tatggaaaat tgatttttta agtaaagtta taacttttaa 3360
141 aattgtatcc tacatctaac tccaaataaa ggtttaaaaa caactatgag caatataagt 3420
143 aatacattta aaatacatat aagagaaaga taaggaaaaa aggaatgact catgaagggt 3480
145 agtacacaat ctatgcattc tgaatatatt cacacttacc aagtatttgg ctccagggtt 3540
147 tctggcagct aatgcaaaga gaggaacaga atcaagtttc atggtattat ctggtagact 3600
149 gtggaagcta tagcatttct gccccctcat gttttcacat tcccccttag agaacagcac 3664
151 aata
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155 <211> LENGTH: 22
156 <212> TYPE: DNA
157 <213> ORGANISM: Artificial Sequence
159 <220> FEATURE:
160 <223> OTHER INFORMATION: synthetic primer
162 <400> SEQUENCE: 2
163 actaagaagt gcattagtcg gg
166 <210> SEQ ID NO: 3
167 <211> LENGTH: 20

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168 <212> TYPE: DNA
169 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: synthetic primer
174 <400> SEQUENCE: 3
175 ttccctgtgct ctagcttgct 20
178 <210> SEQ ID NO: 4
179 <211> LENGTH: 20
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: synthetic primer
186 <400> SEQUENCE: 4
187 tgcaaatacta tgctgcaaaa 20
190 <210> SEQ ID NO: 5
191 <211> LENGTH: 20
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:
196 <223> OTHER INFORMATION: synthetic primer
198 <400> SEQUENCE: 5
199 gggtgcctaa tcacgagaaa 20
202 <210> SEQ ID NO: 6
203 <211> LENGTH: 25
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: synthetic primer
210 <400> SEQUENCE: 6
211 ccaaaggctt ggtgatttag tggac 25
214 <210> SEQ ID NO: 7
215 <211> LENGTH: 25
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: synthetic primer
222 <400> SEQUENCE: 7
223 ctagattgaa ggccagaaaa catgc 25
226 <210> SEQ ID NO: 8
227 <211> LENGTH: 19
228 <212> TYPE: DNA
229 <213> ORGANISM: Artificial Sequence
231 <220> FEATURE:
232 <223> OTHER INFORMATION: synthetic primer
234 <400> SEQUENCE: 8
235 aacatcttag ggcatacctg 19
238 <210> SEQ ID NO: 9
239 <211> LENGTH: 25
240 <212> TYPE: DNA

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241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <223> OTHER INFORMATION: synthetic primer
246 <400> SEQUENCE: 9
247 aatgatttaa aatagattag gagca
250 <210> SEQ ID NO: 10
251 <211> LENGTH: 62
252 <212> TYPE: PRT
253 <213> ORGANISM: Homo sapiens
255 <400> SEQUENCE: 10
257 Met Val Arg Ser Gln Val Glu Trp Lys Gly Gln Leu Ile Pro Ala Ala
258 1 5 10 15
261 Gly Ser Ala Cys Thr His Met Pro Pro Phe Ser Cys Leu Leu Thr Gly
262 20 25 30
265 Ser Ile Glu Gly Val His Asn Glu Ala Ser Cys Lys Thr Ser Pro Asn
266 35 40 45
269 Ser Arg Arg Ser Arg Asn Thr Leu His Leu Gln Arg Asn Leu
270 50 55 60
273 <210> SEQ ID NO: 11
274 <211> LENGTH: 45
275 <212> TYPE: PRT
276 <213> ORGANISM: Homo sapiens
278 <400> SEQUENCE: 11
280 Met Val Arg Ser Gln Val Glu Trp Lys Gly Gln Leu Ile Pro Ala Ala
281 1 5 10 15
284 Gly Ser Ala Cys Thr His Met Pro Pro Phe Ser Cys Leu Leu Thr Gly
285 20 25 30
288 Ser Ile Glu Gly Val His Asn Glu Ala Arg Asp Gly Pro
289 35 40 45
292 <210> SEQ ID NO: 12
293 <211> LENGTH: 189
294 <212> TYPE: DNA
295 <213> ORGANISM: Homo sapiens
297 <400> SEQUENCE: 12
298 atggtgagga gccaaagtga atggaaagga cagctcatcc cggcggctgg gagtgcacgc 60
300 acacacatgc cccctttttc ttgcctacta acaggatcta tagaaggcgt acataatgaa 120
302 gcaagttgca agacaagtcc aaattcaaga aggtcaagaa atacactcca tctccagaga 180
304 aatctatga 189
307 <210> SEQ ID NO: 13
308 <211> LENGTH: 138
309 <212> TYPE: DNA
310 <213> ORGANISM: Homo sapiens
312 <400> SEQUENCE: 13
313 atggtgagga gccaaagtga atggaaagga cagctcatcc cggcggctgg gagtgcacgc 60
315 acacacatgc cccctttttc ttgcctacta acaggatcta tagaaggcgt acataatgaa 120
317 gcaagagatg gaccctga 138
320 <210> SEQ ID NO: 14
321 <211> LENGTH: 20
322 <212> TYPE: DNA

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325 <220> FEATURE:
326 <223> OTHER INFORMATION: synthetic primer
328 <400> SEQUENCE: 14
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332 <210> SEQ ID NO: 15
333 <211> LENGTH: 21
334 <212> TYPE: DNA
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: synthetic primer
340 <400> SEQUENCE: 15
341 ctcccttcac cacaattggt c                                21
344 <210> SEQ ID NO: 16
345 <211> LENGTH: 21
346 <212> TYPE: DNA
347 <213> ORGANISM: Artificial Sequence
349 <220> FEATURE:
350 <223> OTHER INFORMATION: synthetic primer
352 <400> SEQUENCE: 16
353 tcacgcgatta gccacctctt c                                21
356 <210> SEQ ID NO: 17
357 <211> LENGTH: 21
358 <212> TYPE: DNA
359 <213> ORGANISM: Artificial Sequence
361 <220> FEATURE:
362 <223> OTHER INFORMATION: synthetic primer
364 <400> SEQUENCE: 17
365 tgtcaagcac aatgctaattg c                                21
368 <210> SEQ ID NO: 18
369 <211> LENGTH: 23
370 <212> TYPE: DNA
371 <213> ORGANISM: Artificial Sequence
373 <220> FEATURE:
374 <223> OTHER INFORMATION: synthetic primer
376 <400> SEQUENCE: 18
377 gggttgatac cagagtgttc tcc                                23
380 <210> SEQ ID NO: 19
381 <211> LENGTH: 21
382 <212> TYPE: DNA
383 <213> ORGANISM: Artificial Sequence
385 <220> FEATURE:
386 <223> OTHER INFORMATION: synthetic primer
388 <400> SEQUENCE: 19
389 gtcttatgac cagccagcaa t                                21
392 <210> SEQ ID NO: 20
393 <211> LENGTH: 21
394 <212> TYPE: DNA
395 <213> ORGANISM: Artificial Sequence

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/578,701

DATE: 05/18/2006

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Input Set : A:\Sequence Listing.txt

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date